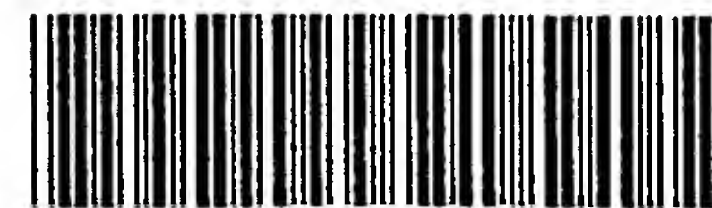


RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/582,606
Source: IFWP
Date Processed by STIC: 8/8/06

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 08/08/2006

PATENT APPLICATION: US/10/587,606

TIME: 08:59:59

Input Set : A:\68074-A-PCT_SEQ_LIST.ST25.txt

Output Set: N:\CRF4\08082006\J587606.raw

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3 <110> APPLICANT: Fine, Robert L
4      Brandt-Rauf, Paul
5      Mao, Yueha
7 <120> TITLE OF INVENTION: C-Terminal p53 Palindromic Peptide That Induces Apoptosis of
8      Cells with Aberrant p53 and Uses Thereof
10 <130> FILE REFERENCE: 68074-A-PCT-US/JPW/JW
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/587,606
C--> 12 <141> CURRENT FILING DATE: 2006-07-27
12 <150> PRIOR APPLICATION NUMBER: PCT/US2005/002543
13 <151> PRIOR FILING DATE: 2005-01-27
15 <150> PRIOR APPLICATION NUMBER: US 60/540,864
16 <151> PRIOR FILING DATE: 2004-01-30
18 <160> NUMBER OF SEQ ID NOS: 9
20 <170> SOFTWARE: PatentIn version 3.3
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 41
24 <212> TYPE: PRT
25 <213> ORGANISM: Artificial Sequence
27 <220> FEATURE:
28 <223> OTHER INFORMATION: Polypeptide based on human p53
30 <400> SEQUENCE: 1
32 Ala Gln Ala Gly Lys Glu Pro Gly Gly Ser Arg Ala His Ser Ser His
33 1          5          10          15
36 Leu Lys Ser Lys Lys Gly Gln Ser Thr Ser Arg His Lys Lys Leu Met
37          20          25          30
40 Phe Lys Thr Glu Gly Pro Asp Ser Asp
41          35          40
44 <210> SEQ ID NO: 2
45 <211> LENGTH: 41
46 <212> TYPE: PRT
47 <213> ORGANISM: Artificial Sequence
49 <220> FEATURE:
50 <223> OTHER INFORMATION: Polypeptide based on human p53
52 <400> SEQUENCE: 2
54 Asp Ser Asp Pro Gly Glu Thr Lys Phe Met Leu Lys Lys His Arg Ser
55 1          5          10          15
58 Thr Ser Gln Gly Lys Lys Ser Lys Leu His Ser Ser His Ala Arg Ser
59          20          25          30
62 Gly Gly Pro Glu Lys Gly Ala Gln Ala
63          35          40
66 <210> SEQ ID NO: 3
67 <211> LENGTH: 83
68 <212> TYPE: PRT

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RAW SEQUENCE LISTING

DATE: 08/08/2006

PATENT APPLICATION: US/10/587,606

TIME: 08:59:59

Input Set : A:\68074-A-PCT_SEQ_LIST.ST25.txt

Output Set: N:\CRF4\08082006\J587606.raw

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69 <213> ORGANISM: Artificial Sequence
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72 <223> OTHER INFORMATION: Palindromic polypeptide based on human p53
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76 <221> NAME/KEY: MISC_FEATURE
77 <222> LOCATION: (42)..(42)
78 <223> OTHER INFORMATION: Xaa = Gly or absent
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83 1 5 10 15
86 Leu Lys Ser Lys Lys Gly Gln Ser Thr Ser Arg His Lys Lys Leu Met
87 20 25 30
W--> 90 Phe Lys Thr Glu Gly Pro Asp Ser Asp Xaa Asp Ser Asp Pro Gly Glu
91 35 40 45
94 Thr Lys Phe Met Leu Lys Lys His Arg Ser Thr Ser Gln Gly Lys Lys
95 50 55 60
98 Ser Lys Leu His Ser Ser His Ala Arg Ser Gly Gly Pro Glu Lys Gly
99 65 70 75 80
102 Ala Gln Ala
106 <210> SEQ ID NO: 4
107 <211> LENGTH: 83
108 <212> TYPE: PRT
109 <213> ORGANISM: Artificial Sequence
111 <220> FEATURE:
112 <223> OTHER INFORMATION: Palindromic polypeptide based on human p53
115 <220> FEATURE:
116 <221> NAME/KEY: MISC_FEATURE
117 <222> LOCATION: (42)..(42)
118 <223> OTHER INFORMATION: Xaa = Gly or absent
120 <400> SEQUENCE: 4
122 Asp Ser Asp Pro Gly Glu Thr Lys Phe Met Leu Lys Lys His Arg Ser
123 1 5 10 15
126 Thr Ser Gln Gly Lys Lys Ser Lys Leu His Ser Ser His Ala Arg Ser
127 20 25 30
W--> 130 Gly Gly Pro Glu Lys Gly Ala Gln Ala Xaa Ala Gln Ala Gly Lys Glu
131 35 40 45
134 Pro Gly Gly Ser Arg Ala His Ser Ser His Leu Lys Ser Lys Lys Gly
135 50 55 60
138 Gln Ser Thr Ser Arg His Lys Lys Leu Met Phe Lys Thr Glu Gly Pro
139 65 70 75 80
142 Asp Ser Asp
146 <210> SEQ ID NO: 5
147 <211> LENGTH: 167
148 <212> TYPE: PRT
149 <213> ORGANISM: Artificial Sequence
151 <220> FEATURE:
152 <223> OTHER INFORMATION: Palindromic polypeptide based on human p53
155 <220> FEATURE:
156 <221> NAME/KEY: MISC_FEATURE

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RAW SEQUENCE LISTING

DATE: 08/08/2006

PATENT APPLICATION: US/10/587,606

TIME: 08:59:59

Input Set : A:\68074-A-PCT_SEQ_LIST.ST25.txt

Output Set: N:\CRF4\08082006\J587606.raw

157 <222> LOCATION: (42)..(42)
 158 <223> OTHER INFORMATION: Xaa = Gly or absent
 160 <220> FEATURE:
 161 <221> NAME/KEY: MISC_FEATURE
 162 <222> LOCATION: (84)..(84)
 163 <223> OTHER INFORMATION: Xaa = Gly or absent
 165 <220> FEATURE:
 166 <221> NAME/KEY: MISC_FEATURE
 167 <222> LOCATION: (126)..(126)
 168 <223> OTHER INFORMATION: Xaa = Gly or absent
 170 <400> SEQUENCE: 5
 172 Ala Gln Ala Gly Lys Glu Pro Gly Gly Ser Arg Ala His Ser Ser His
 173 1 5 10 15
 176 Leu Lys Ser Lys Lys Gly Gln Ser Thr Ser Arg His Lys Lys Leu Met
 177 20 25 30
 W--> 180 Phe Lys Thr Glu Gly Pro Asp Ser Asp Xaa Asp Ser Asp Pro Gly Glu
 181 35 40 45
 184 Thr Lys Phe Met Leu Lys Lys His Arg Ser Thr Ser Gln Gly Lys Lys
 185 50 55 60
 188 Ser Lys Leu His Ser Ser His Ala Arg Ser Gly Gly Pro Glu Lys Gly
 189 65 70 75 80
 192 Ala Gln Ala Xaa Ala Gln Ala Gly Lys Glu Pro Gly Gly Ser Arg Ala
 193 85 90 95
 196 His Ser Ser His Leu Lys Ser Lys Lys Gly Gln Ser Thr Ser Arg His
 197 100 105 110
 200 Lys Lys Leu Met Phe Lys Thr Glu Gly Pro Asp Ser Asp Xaa Asp Ser
 201 115 120 125
 204 Asp Pro Gly Glu Thr Lys Phe Met Leu Lys Lys His Arg Ser Thr Ser
 205 130 135 140
 208 Gln Gly Lys Lys Ser Lys Leu His Ser Ser His Ala Arg Ser Gly Gly
 209 145 150 155 160
 212 Pro Glu Lys Gly Ala Gln Ala
 213 165
 216 <210> SEQ ID NO: 6
 217 <211> LENGTH: 167
 218 <212> TYPE: PRT
 219 <213> ORGANISM: Artificial Sequence
 221 <220> FEATURE:
 222 <223> OTHER INFORMATION: Palindromic polypeptide based on human p53
 225 <220> FEATURE:
 226 <221> NAME/KEY: MISC_FEATURE
 227 <222> LOCATION: (42)..(42)
 228 <223> OTHER INFORMATION: Xaa = Gly or absent
 230 <220> FEATURE:
 231 <221> NAME/KEY: MISC_FEATURE
 232 <222> LOCATION: (84)..(84)
 233 <223> OTHER INFORMATION: Xaa = Gly or absent
 235 <220> FEATURE:
 236 <221> NAME/KEY: MISC_FEATURE

RAW SEQUENCE LISTING

DATE: 08/08/2006

PATENT APPLICATION: US/10/587,606

TIME: 08:59:59

Input Set : A:\68074-A-PCT_SEQ_LIST.ST25.txt

Output Set: N:\CRF4\08082006\J587606.raw

237 <222> LOCATION: (126)..(126)

238 <223> OTHER INFORMATION: Xaa = Gly or absent

240 <400> SEQUENCE: 6

242	Asp	Ser	Asp	Pro	Gly	Glu	Thr	Lys	Phe	Met	Leu	Lys	Lys	His	Arg	Ser
243	1				5				10					15		
246	Thr	Ser	Gln	Gly	Lys	Lys	Ser	Lys	Leu	His	Ser	Ser	His	Ala	Arg	Ser
247				20					25					30		
W--> 250	Gly	Gly	Pro	Glu	Lys	Gly	Ala	Gln	Ala	Xaa	Ala	Gln	Ala	Gly	Lys	Glu
251				35				40					45			
254	Pro	Gly	Gly	Ser	Arg	Ala	His	Ser	Ser	His	Leu	Lys	Ser	Lys	Lys	Gly
255		50					55					60				
258	Gln	Ser	Thr	Ser	Arg	His	Lys	Lys	Leu	Met	Phe	Lys	Thr	Glu	Gly	Pro
259	65					70					75					80
262	Asp	Ser	Asp	Xaa	Asp	Ser	Asp	Pro	Gly	Glu	Thr	Lys	Phe	Met	Leu	Lys
263				85					90						95	
266	Lys	His	Arg	Ser	Thr	Ser	Gln	Gly	Lys	Lys	Ser	Lys	Leu	His	Ser	Ser
267				100					105					110		
270	His	Ala	Arg	Ser	Gly	Gly	Pro	Glu	Lys	Gly	Ala	Gln	Ala	Xaa	Ala	Gln
271				115					120					125		
274	Ala	Gly	Lys	Glu	Pro	Gly	Gly	Ser	Arg	Ala	His	Ser	Ser	His	Leu	Lys
275		130					135						140			
278	Ser	Lys	Lys	Gly	Gln	Ser	Thr	Ser	Arg	His	Lys	Lys	Leu	Met	Phe	Lys
279	145					150					155					160
282	Thr	Glu	Gly	Pro	Asp	Ser	Asp									
283						165										

286 <210> SEQ ID NO: 7

287 <211> LENGTH: 164

288 <212> TYPE: PRT

289 <213> ORGANISM: Artificial Sequence

291 <220> FEATURE:

292 <223> OTHER INFORMATION: Palindromic polypeptide based on human p53

294 <400> SEQUENCE: 7

296	Ala	Gln	Ala	Gly	Lys	Glu	Pro	Gly	Gly	Ser	Arg	Ala	His	Ser	Ser	His
297	1				5				10					15		
300	Leu	Lys	Ser	Lys	Lys	Gly	Gln	Ser	Thr	Ser	Arg	His	Lys	Lys	Leu	Met
301				20					25					30		
304	Phe	Lys	Thr	Glu	Gly	Pro	Asp	Ser	Asp	Asp	Ser	Asp	Pro	Gly	Glu	Thr
305			35					40					45			
308	Lys	Phe	Met	Leu	Lys	Lys	His	Arg	Ser	Thr	Ser	Gln	Gly	Lys	Lys	Ser
309		50					55					60				
312	Lys	Leu	His	Ser	Ser	His	Ala	Arg	Ser	Gly	Gly	Pro	Glu	Lys	Gly	Ala
313	65					70					75					80
316	Gln	Ala	Ala	Gln	Ala	Gly	Lys	Glu	Pro	Gly	Gly	Ser	Arg	Ala	His	Ser
317				85					90					95		
320	Ser	His	Leu	Lys	Ser	Lys	Lys	Gly	Gln	Ser	Thr	Ser	Arg	His	Lys	Lys
321				100					105					110		
324	Leu	Met	Phe	Lys	Thr	Glu	Gly	Pro	Asp	Ser	Asp	Asp	Ser	Asp	Pro	Gly
325			115					120					125			
328	Glu	Thr	Lys	Phe	Met	Leu	Lys	Lys	His	Arg	Ser	Thr	Ser	Gln	Gly	Lys

RAW SEQUENCE LISTING

DATE: 08/08/2006

PATENT APPLICATION: US/10/587,606

TIME: 08:59:59

Input Set : A:\68074-A-PCT_SEQ_LIST.ST25.txt

Output Set: N:\CRF4\08082006\J587606.raw

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329      130      135      140
332 Lys Ser Lys Leu His Ser Ser His Ala Arg Ser Gly Gly Pro Glu Lys
333 145      150      155      160
336 Gly Ala Gln Ala
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341 <211> LENGTH: 17
342 <212> TYPE: PRT
343 <213> ORGANISM: Artificial Sequence
345 <220> FEATURE:
346 <223> OTHER INFORMATION: Membrane carrier peptide derived from Antennaepedia
348 <400> SEQUENCE: 8
350 Lys Lys Trp Lys Met Arg Arg Asn Gln Phe Trp Val Lys Val Gln Arg
351 1      5      10      15
354 Gly
358 <210> SEQ ID NO: 9
359 <211> LENGTH: 6
360 <212> TYPE: DNA
361 <213> ORGANISM: Artificial Sequence
363 <220> FEATURE:
364 <223> OTHER INFORMATION: Restriction enzyme site derived from human p53
366 <400> SEQUENCE: 9
367 ggccgg

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6

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/08/2006
PATENT APPLICATION: US/10/587,606 TIME: 09:00:00

Input Set : A:\68074-A-PCT_SEQ_LIST.ST25.txt
Output Set: N:\CRF4\08082006\J587606.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 42 '
Seq#:4; Xaa Pos. 42
Seq#:5; Xaa Pos. 42, 84, 126
Seq#:6; Xaa Pos. 42, 84, 126

VERIFICATION SUMMARY

DATE: 08/08/2006

PATENT APPLICATION: US/10/587,606

TIME: 09:00:00

Input Set : A:\68074-A-PCT_SEQ_LIST.ST25.txt

Output Set: N:\CRF4\08082006\J587606.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:90 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:32
L:130 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:32
L:180 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:32
M:341 Repeated in SeqNo=5
L:250 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:32
M:341 Repeated in SeqNo=6